

Figure 2

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G(olf) Galphas XLG(olf) XLGalphas	
G(olf) Galphas XLG(olf) XLGalphas	MGCLGGNSKTTBDQGVDEKEMGCLGGNSKTTBDQGVDEKE
G(olf) Galphas XLG(olf) XLGalphas	βγ subunit binding RREANKKIEKQLQKERLAYKATHRLLLLGAGESGKSTIVKQMRILHVNGFNPEE QREANKKIEKQLQKDKQVYRATHRLLLLGAGESGKSTIVKQMRILHVNGFNGEGEEDPQ ARKVSRGIDRMLRDQKRDLQQTHRLLLLGAGESGKSTIVKQMRILHVNGFNPEE EKKRSKLIDKQLQDEKMGYMCTHRLLLLGAGESGKSTIVKQMRILHVNGFNGEGGEEDPQ EKKRSKLIDKQLQDEKMGYMCTHRLLLLGAGESGKSTIVKQMRILHVNGFNGEGGEEDPQ
G(olf) Galphas XLG(olf) XLGalphas	I KKQKILDIRKNVKDAIVŢIVSAMSTIIPPVPLANPENQFRSDYIKSIAPIT AARSNSDGEKATKVQDIKNNLKEAIETIVAAMSNLVPPVELANPENQFRVDYILSVMNVP KKQKILDIRKNVKDAIVŢIVSAMSTIIPPVPLANPENQFRSDYIKSIAPIT AARSNSDGEKAŢKVQDIKNNLKEAIETIVAAMSNLVPPVELANPENQFRVDYILSVMNVP

Figure 4

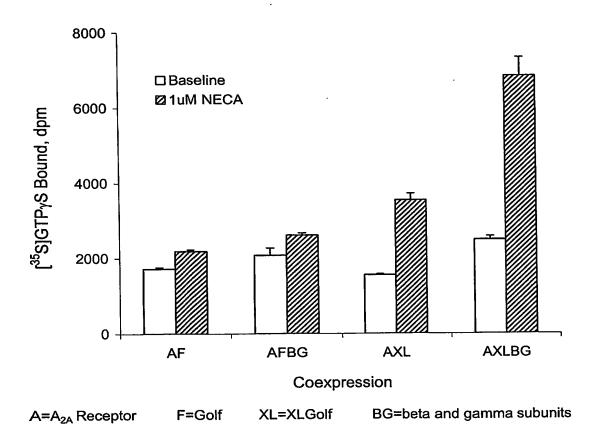


Figure 5

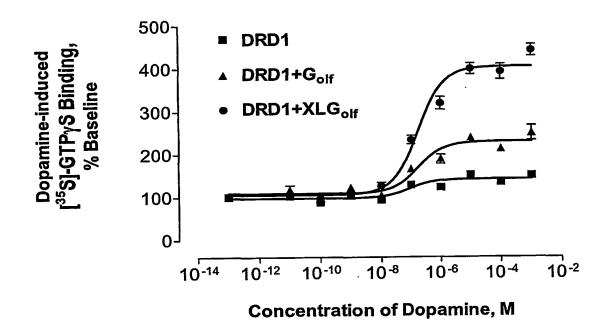


Figure 6

T98G Glioma DNA Golf and XL Golf MSP

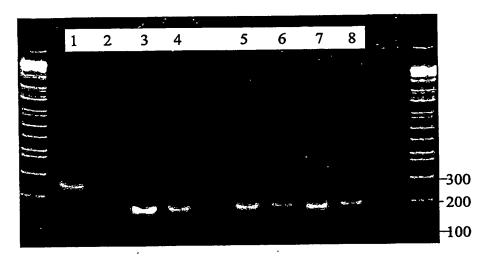
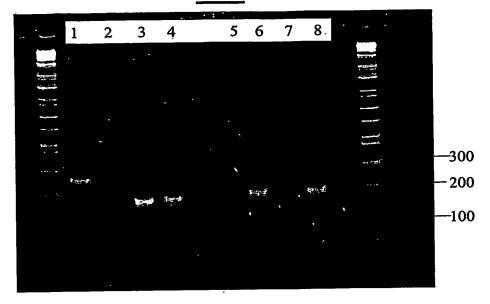


Figure 7

Peripheral Blood DNA Golf and XL Golf MSP



WO 2005/047318 PCT/GB2004/004749

Figure 8

Human GNAL locus; 197140 nt

Features	
Alternate exon 1	20012685
CpG island	21182881
Variation T/C	2198
Variation A/G	2216
Variation C/T	2268
XLGolf start codon	2309
Variation repeat	24212426
Variation A/G	2445
Variation repeat	27462765
Variation C/T	2765
CpG island	6425365647
Variation T/G	64734
Variation C/T	64748
Exon 1	6492565323
Golf start codon	65179
Exon 2	6559865670
Exon 3	6637366427
Exon 4	6657166690
Exon 5	137663137760
Exon 6	175140175194
Exon 7	177278177351
Exon 8	179913179971
Exon 9	181288181408
Exon 10	185013185143
Exon 11	189366189433
Exon 12	193734194571
Stop codon	193878

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Fig. 8 (cont.)						
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Dennassesn	taacaaaata	cagagatete	tgcaagaaat	ggattgggga	aattgaagge	1680
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Fig. 8 (cont.)

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Fig. 8 (cont.)						
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Fig. 8 (cont.)

g. 8 (cont.)						
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Fig. 8 (cont.)						
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Fig. 8 (cont.)						
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Fig. 8 (cont.)

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Fig. 8 (cont.)

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gcacggccac	caataacaac	aaacattact	gctacccgca	cttcacctgc	geegragaea	193800
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catetateca	: attctatatc	gaccaccaag	cctctggcta	cctctgtccc	ctcaggtttg	194040
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ctggtttatt	: acactgtaca	gaccacaaaa	tgtaatattc	: ttttgtataa	ctactaaaga	
aaaatcctto	, tagatctttg	tgccttcacc	atggctatct	. atacctgtac	atgaaatgtg	194700 194760
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ctgatttaaa	a aaattgtaag	ttatacacgt	. taatcatcca	cattctatcg	acaatgtacc	194940
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caattcaggg	g gtacaaattg	caatctaatc	: ttttcaggga	accagggatt	tttttctctc	195060
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g. o (conc.)						
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totataaaga	tatccaaaat	taaattttca	acaatacaaa	tctagagaag	tgacagccta	195780
cattacttca	trattactct	tettttagte	tttagtcttt	aatatttaa	agtttactct	195840
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acaaaccage	teactttata	gacagtcagt	gcaacacaca	cattttatct	catcaccgtc	195960
ttaataasta	cccatccact	gtctcataaa	gccctcagct	gaactaagat	aaataataca	196020
ntacigotto	ttttcacttc	cccttacact	gtcaaagtaa	aacaagaaaa	ctgaaaagct	196080
acygaaacca	caadaaadd	aagtatgctg	ttgtatgcat	cattactcaa	caattaccct	196140
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ceteagetgg	taccacagge	accedence	gtcttgaact	cctgacctca	agtgatccac	196680
agacgaggii	ctaccatget	actaggets	caggtgtgag	ccactgtgcc	cggccacatc	196740
cegeeteage	aggetagage	tttacactac	ataggagattt	gagaaccact	tcacaggaag	196800
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agggaaacag	cccaatattt	caaggactat	catacatata	agtgacgaca	ttaatagcat	196920
ceaggeeett	trangatga	caaggactgt	atacatatat	ttgataaaaa	tgagaaaaca	196980
ttacatactg	racayaryca	togactttta	taggatgaga	acagtacaat	caactattta	197040
gatttgttgt	agageacceg	atasttasas	atttctatac	tgtgcagaga	gacggcctgt	197100
ttttgcagtt	accoactica	tgattctaac	atrateteto	cycycagaga	5555-	197140
aattggtctc	atcatccact	Lyallulado	acyacticity			

Figure 9

Amino Acid Sequence of Human Golf

1	MGCLGGNSKT	TEDOGVDEKE	RREANKKIEK	OLOKEKLYAK	ATHRLLLLGA	GESGKSTIVK
61	OMRILHVNGF	NPEEKKOKIL	DIRKNVKDAI	VTIVSAMSTI	IPPVPLANPE	NQFRSDYIKS
121	TAPITDFEYS	OEFFDHVKKL	WDDEGVKACF	ERSNEYQLID	CAQYFLERID	SVSLVDYTPT
181	DODLIBCRVL	TSGIFETRFO	VDKVNFHMFD	VGGQRDERIK	WIQCFNDVTA	IIYVAACSSY
241	NMVTREDNNT	NRLRESLDLF	ESIWNNRWLR	TISIILFLNK	QDMLAEKVLA	GKSKIEDYFP
301	EVANYTVPED	ATPDAGEDPK	VTRAKFFIRD	LFLRISTATG	DGKHYCYPHF	TCAVDTENIR
	RVFNDCRDII					

Figure 10

cDNA and Amino Acid Sequence of Human XLGolf

	MGLCYSLRPLLFGGPGDDPC
1	ATGGGTCTGTGCTACAGTCTGCGGCCGCTGCTTTTCGGGGGGCCCAGGGGACGACCCCTGC
	AASEPPVEDAQPAPALAP
61	GCGGCCTCGGAGCCGCGGTGGAGGACGCGCAGCCCGGCCCCGGCCCCGGCCCCCA
	V R A A A R D T A R T L L P R G G E G S
121	GTCCGGGCGGCCGCAAGGGACACGGCCCGGACCCTGCTCCCTCGGGGCGGCGAAGGGAGC
	PACARPKADKPKEKRQRTEQ
181	CCGGCATGCGCTCGGCCCAAAGCAGACAAGCCGAAGGAGAAGCGGCAGCGCACCGAGCAG
	LSAEEREAAKEREAVKEARK
241	CTGAGTGCCGAGGAGCGCGAGGCGCCAAGGAGCGCGAGGCGAGGAAA
	V S R G I D R M L R D Q K R D L Q Q T H
301	GTGAGCCGGGGCATCGACCGCATGCTGCGCGACCAGAAGCGCGACCTGCAGCAGACGCAC
	R. L L L G A G E S G K S T I V K Q M R
361	CGGCTCCTGCTCGGGGCTGGTGAGTCTGGGAAAAGCACCATCGTCAAACAGATGAGG
	I. L H V N G F N P E E K K Q K I L D I R ATCCTGCACGTCAATGGGTTTAATCCCGAGGAAAAGAAACAGAAAATTCTGGACATCCGG
421	
401	K N V K D A I V T I V S A M S T I I P P AAAAATGTTAAAGATGCTATCGTGACAATTGTTTCAGCAATGAGTACTATAATACCTCCA
481	V P L A N P E N Q F R S D Y I K S I A P
541	GTTCCGCTGGCCAACCCTGAAAACCAATTTCGATCAGACTACATCAAGAGCATAGCCCCT
241	I T D F E Y S Q E F F D H V K K L W D D
601	ATCACTGACTTTGAATATTCCCAGGAATTCTTTGACCATGTGAAAAAACTTTGGGACGAT
001	E.G.V.K.A.C.F.E.R.S.N.E.Y.Q.L.I.D.C.A.Q
661	GAAGGCGTGAAGGCATGCTTTGAGAGATCCAACGAATACCAGCTGATTGACTGTGCACAA
001	YFLERIDSVSLVDYTPTDQD
721	TACTTCCTGGAAAGAATCGACAGCGTCAGCTTGGTTGACTACACACCCCACAGACCAGGAC
	LLRCRVLTSGIFETRFQVDK
781	CTCCTCAGATGCAGAGTTCTGACATCTGGGATTTTTGAGACACGATTCCAAGTGGACAAA
	VNFHMFDVGGQRDERRKWIQ
841	GTAAACTTCCACATGTTTGATGTTGGTGGCCAGAGGGATGAGAGGAGAAAATGGATCCAG
	C F N D V T A I I Y V A A C S S Y N M V
901	TGCTTTAACGATGTCACAGCTATCATTTACGTCGCAGCCTGCAGTAGCTACAACATGGTG
	I R. E D N N T N R L R E S L D L F E S I
961	ATTCGAGAAGATAACAACACCAACAGGCTGAGAGAGTCCCTGGATCTTTTTGAAAGCATC
	M M M M M T K T T D T T D T T D T T T
1021	TGGAACACAGGTGGTTACGGACCATTTCTATCATCTTGTTCTTGAACAAACA
	L A E K V L A G K S K I E D Y F P E Y A CTGGCAGAAAAAGTCTTGGCAGGGAAATCAAAAATTGAAGACTATTTCCCAGAATATGCA
1081	
1111	N Y T V P E D A T P D A G E D P K V T R AATTATACTGTTCCTGAAGACGCAACACCAGATGCAGGAAGATCCCAAAGTTACAAGA
1141	A K F F I R D L F L R I S T A T G D G K
1201	GCCAAGTTCTTTATCCGGGACCTGTTTTTGAGGATCAGCACGGCCACCGGTGACGGCAAA
1201	H Y C Y P H F T C A V D T E N I R R V F
1261	CATTACTGCTACCCGCACTTCACCTGCGCCGTGGACACAGAGAACATCCGCAGGGTGTTC
1201	N D C R D I I Q R M H L K Q Y E L L *
1321	AACGACTGCCGCGACATCATCCAGCGGATGCACCTCAAGCAGTATGAGCTCTTGTGA

Figure 11

G _{olf} G _{as} XLG _{olf} XLG _{as}	MEISGPPFEIGSAPAGVDDTPVNMDSPPIALDGPPIKVSGAPDKRERAERPPVEEEAAEM
G _{olf} G _{as} XLG _{olf} XLG _{as}	EGAADAAEGGKVPSPGYGSPAAGAASADTAARAAPAAPADPDSGATPEDPDSGTAPADPD
G _{olf} G _{αs} XLG _{olf} XLG _{αs}	SGAFAADPDSGAAPAAPADPDSGAAPDAPADPDAGAA PEAPAAPAAAE
G _{olf} G _{as} XLG _{olf} XLG _{as}	TRAAHVAPAAPDAGAPTAPAASATRAAQVRRAASAAPASGARRKIHLRPPSPEIQAADPP
G _{olf} G _{αs} XLG _{olf} XLG _{αs}	TPRPTRASAWRGKSESSRGRRVYYDEGVASSDDDSSGDESDDGTSGCLRWFQHRRNRRRR
G _{olf} G _{os} XLG _{olf} XLG _{os}	LAPVRAAARDTARTLLPRGGEGSPACARPKADKPKEKRQRTEQLSAEER EAAKEREAVKE KPQRNLLRNFLVQAFGGCFGRSESPQPKASRSLKVKKVPLAEKRQMRK EALEKRAQKRA
G _{olf} G _{as} XLG _{olf} XLG _{as}	βγ binding domain RREANKKIEKQLQKERLAYKATHRLLLLGAGESGKSTIVKQMRILHVNGFNPE QREANKKIEKQLQKDKQVYRATHRLLLLGAGESGKSTIVKQMRILHVNGFNGE ARKVSRGIDRMLRDQKRDLQQTHRLLLLGAGESGKSTIVKQMRILHVNGFNPE EKKRSKLIDKQLQDEKMGYMCTHRLLLLGAGESGKSTIVKQMRILHVNGFNGE

Figure 12

	1				50
G(olf) XL-G(olf)				cttttcgggg	
C/c15)	51		~~~~~~~		100
G(olf)	cgacccctgc	geggeetegg	agccgccggt	ggaggacgcg	cagcccgccc
G(olf)	101	~~~~~~	~~~~~~		150
XL-G(olf)	eggeceegge	cctggcccca	gtccgggcgg	ccgcaaggga	cacggcccgg
G(olf)	151			~~~~~~	200
XL-G(olf)	accetgetee	ctcggggcgg	cgaagggagc	ccggcatgcg	ctcggcccaa
G(olf) XL-G(olf)	201 ~~~~~~ agcagacaag	 ccgaaggaga	ageggeageg	~ <u>AtaGaGtat</u> cAccGaGcag	250 <u>tTGqGcGqCa</u> cTGaGtGcCg
G(olf) XL-G(olf)	251 <u>Acaqcaaqac</u> Aggagcgcga	GaCGGaagAc GgCGGccaAg	cAGgGCGtcG gAGcGCGagG	atGaaAAaGA cgGtcAAgGA	300 <u>acqacGcqAq</u> ggcgaGgaAa
G(olf) XL-G(olf)	301 <u>GccAaCaaaa</u> GtgAgCcggg	agATCGAgaa gcATCGAccg	gcaGtTGCag catGcTGCgc	aAaqAGcqcC gAccAGaagC	350 tgGcttacaA gcGacctgcA
G(olf) XL-G(olf)	351 <u>GgctACcCAC</u> GcagACgCAC	CGcCTaCTGC CGgCTcCTGC	TCCTqGGGGC TgCTCGGGGC	TGGTGAGTCT TGGTGAGTCT	400 GGGAAAAGCA GGGAAAAGCA
G(olf) XL-G(olf)	401 CTATCGTCAA CTATCGTCAA	ACAGATGAGG ACAGATGAGG	ATCCTGCACG ATCCTGCACG	TCAATGGGTT TCAATGGGTT	450 TAATCCCGAG TAATCCCGAG
G(olf) XL-G(olf)	451 GAAAAGAAAC GAAAAGAAAC	AGAAAATTCT AGAAAATTCT	GGACATCCGG GGACATCCGG	AAAAATGTTA AAAAATGTTA	500 AAGATGCTAT AAGATGCTAT
G(olf) XL-G(olf)	501 CGTGACAATT CGTGACAATT	GTTTCAGCAA GTTTCAGCAA	TGAGTACTAT TGAGTACTAT	AATACCTCCA AATACCTCCA	550 GTTCCGCTGG GTTCCGCTGG
G(olf) XL-G(olf)	551 CCAACCCTGA CCAACCCTGA	AAACCAATTT AAACCAATTT	CGATCAGACT CGATCAGACT	ACATCAAGAG ACATCAAGAG	600 CATAGCCCCT CATAGCCCCT
G(olf) XL-G(olf)	601 ATCACTGACT ATCACTGACT	TTGAATATTC TTGAATATTC	CCAGGAATTC CCAGGAATTC	TTTGACCATG TTTGACCATG	650 TGAAAAACT TGAAAAAACT
G(olf) XL-G(olf)	651 TTGGGACGAT TTGGGACGAT	GAAGGCGTGA GAAGGCGTGA	AGGCATGCTT AGGCATGCTT	TGAGAGATCC TGAGAGATCC	700 AACGAATACC AACGAATACC
G(olf) XL-G(olf)	701 AGCTGATTGA AGCTGATTGA	CTGTGCACAA CTGTGCACAA	TACTTCCTGG TACTTCCTGG	AAAGAATCGA AAAGAATCGA	750 CAGCGTCAGC CAGCGTCAGC
G(olf) XL-G(olf)	751 TTGGTTGACT TTGGTTGACT	ACACACCCAC ACACACCCAC	AGACCAGGAC AGACCAGGAC	CTCCTCAGAT CTCCTCAGAT	800 GCAGAGTTCT GCAGAGTTCT
G(olf) XL-G(olf)	801 GACATCTGGG GACATCTGGG	ATTTTTGAGA ATTTTTGAGA	CACGATTCCA CACGATTCCA	AGTGGACAAA AGTGGACAAA	850 GTAAACTTCC GTAAACTTCC
G(olf) XL-G(olf)	851 ACATGTTTGA ACATGTTTGA	TGTTGGTGGC TGTTGGTGGC	CAGAGGGATG	AGAGGAGAAA AGAGGAGAAA	900 ATGGATCCAG ATGGATCCAG
G(olf) XL-G(olf)	901 TGCTTTAACG TGCTTTAACG	ATGTCACAGO ATGTCACAGO	TATCATTTAC	GTCGCAGCCT GTCGCAGCCT	950 GCAGTAGCTA GCAGTAGCTA
G(olf) XL-G(olf)	951 CAACATGGTG CAACATGGTG	ATTCGAGAAG ATTCGAGAAG	ATAACAACAC ATAACAACAC	CAACAGGCTG CAACAGGCTG	1000 AGAGAGTCCC AGAGAGTCCC

G(olf) XL-G(olf)	1001 TGGATCTTTT TGGATCTTTT			GGTGGTTACG GGTGGTTACG	
G(olf) XL-G(olf)	1051 ATCATCTTGT ATCATCTTGT	TCTTGAACAA TCTTGAACAA		CTGGCAGAAA CTGGCAGAAA	
G(olf) XL-G(olf)		DAADTTGAAG DAADTTAAAA		AGAATATGCA AGAATATGCA	
G(olf) XL-G(olf)		CGCAACACCA CGCAACACCA		AAGATCCCAA AAGACCCCAA	
G(olf) XL-G(olf)		TTATCCGGGA TTATCCGGGA		AGGATCAGCA AGGATCAGCA	
G(olf) XL-G(olf)		CATTACTGCT CATTACTGCT	ACCCGCACTT ACCCGCACTT	CACCTGCGCC CACCTGCGCC	
G(olf) XL-G(olf)	1301 AGAACATCCG AGAACATCCG	CAGGGTGTTC CAGGGTGTTC	AACGACTGCC AACGACTGCC	GCGACATCAT GCGACATCAT	
G(olf) XL-G(olf)		AGTATGAGCT AGTATGAGCT		•	

Figure 13

	1				50
G(olf)		lfggpgddpc	~~~~~~~~	mananalap	vraaardtar
xl-G(olf)	mgrcAsrrbr	Tiggpgaapc	aaseppveda	dbababa-at	100
0/-16\	51	~~~~~~	~~~~~mac	Lagnsktted	gavdeKErRe
G(olf) $x1-G(olf)$	tllprggegs	pacarpkadk	pkekrartea	Lsaeereaak	ereavKEaRk
	101				150
G(olf)	ankkTekaLa	kerlaykaTH	RLLLLGAGES	GKSTIVKQMR GKSTIVKQMR	ILHVNGFNPE ILHVNGFNPE
xl-G(olf)	vsrgIdrmLr	dqkrdlqqTH	KILLILIGAGES	GKSTIVKQIAK	
	151	KNVKDAIVTI	TICAMOUTTOD	VPLANPENQF	200 RSDYIKSIAP
G(olf) xl-G(olf)	EKKOKILDIR	KNVKDAIVTI	VSAMSTIIPP	VPLANPENQF	RSDYIKSIAP
	201				250
G(olf)	ITDFEYSOEF	FDHVKKLWDD	EGVKACFERS	NEYQLIDCAQ	YFLERIDSVS
x1-G(olf)	ITDFEYSQEF	FDHVKKLWDD	EGVKACFERS	NEYQLIDCAQ	
	251			TATERIMEDIACC	300 QRDERRKWIQ
G(olf) xl-G(olf)	LVDYTPTDQD	LLRCRVLTSG LLRCRVLTSG		VNFHMFDVGG	QRDERRKWIQ
X1-G(011)					350
G(olf)	301 CFNDVTAIIÝ	VAACSSYNMV	IREDNNTNRL	RESLDLFESI	WNNRWLRTIS
x1-G(olf)	CFNDVTAIIY	VAAÇSSYNMV	IREDNNTNRL	RESLDLFESI	WNNRWLRTIS
	351	•			400
G(olf)	IILFLNKQDM	LAEKVLAGKS LAEKVLAGKS	KIEDYFPEYA	NYTVPEDATP	DAGEDPKVTR
x1-G(olf)	•			•	450
G(olf)	401	RISTATGDGK	НҮСҮРНЕТСА	VDTENIRRVF	NDCRDIIQRM
x1-G(olf)	AKFFIRDLFL	RISTATGDGK	HYCYPHFTCA	VDTENIRRVF	NDCRDIIQRM
	451	•			
G(olf)	HLKQYELL				
xl-G(olf)	HLKQYELL				

Figure 14

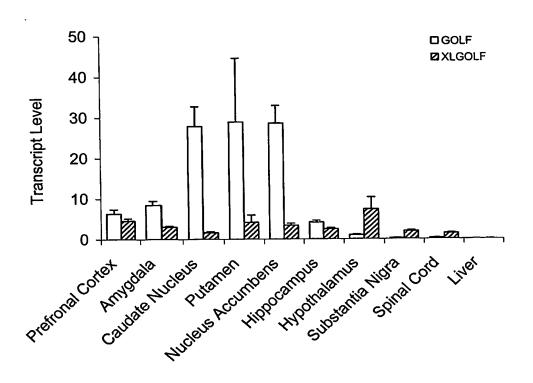


Figure 15

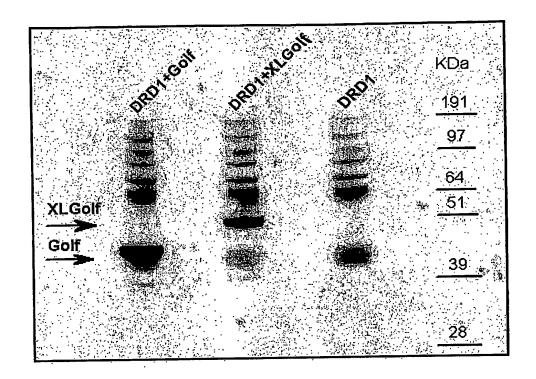
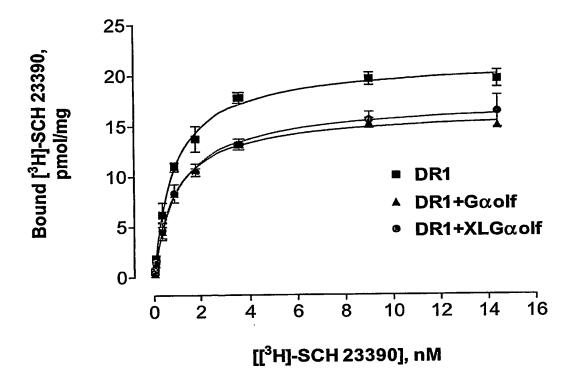
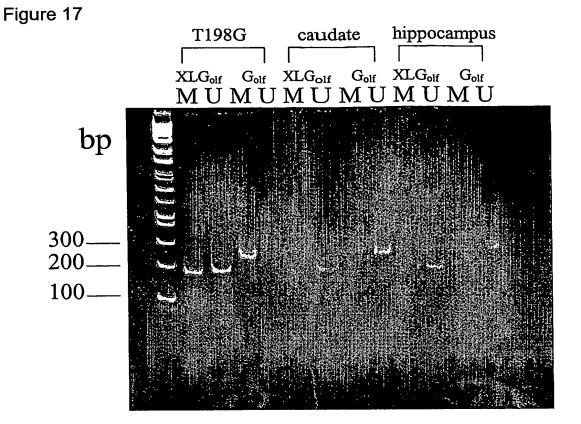
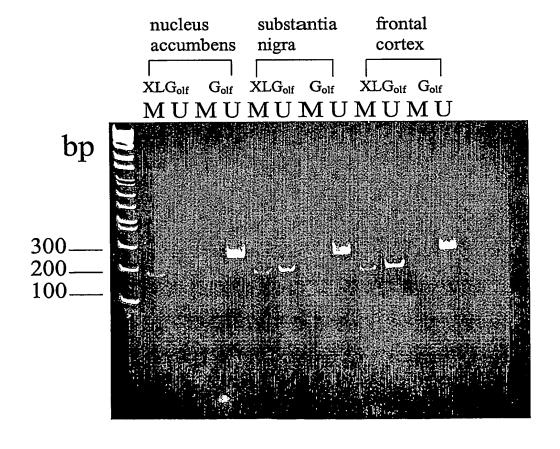


Figure 16







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SUBSTITUTE SHEET (RULE 26)

Figure 18 Mouse XLGolf cDNA

ggcgaggctcgcgcctgggacgttcgaaggatgcacaatggcccttcggggcagtgaggggc cgtggtccccagccggacccaccgcagccaccccgcagctggcgccccagggtccgagt gcagcgctgcctccggcgtcggcggagccccggtctgacccgcggggacttgcccgcgccc cgcgccccgatgggcctatgctacagcctgcggccgctgctcttcgggagcccagaggacac cccgtgtgcggcctcggaaccctgcgcagaggatgctcagcccagcgcccccggcccctg cctcgatcccagccccggctcccgtagggaccctgctccggcgtggcggccggatcgtc gcgaacgcgcggccgccaggcgagctgcagagccgccggcgacaggagcagctacgagccga ggagcgcgaggctaaagaggcgaggaaagtcagccggggcatcgaccgcatgctgcgcg agcagaagcgggacctgcagcagacgcaccggctcctgctgctgggggctggtgagtccggg aaaagcactatcgtcaaacagatgaggatcctgcacgtcaatggcttcaaccccgaggaaaa gaagcagaaaattctggacatcaggaaaaatgtcaaagatgcgatcgtgacaatcgtttcag caatgagtactatcatacctccagttccactggccaaccctgagaaccagttccggtcagat tatatcaagagcatagcccctatcactgactttgaatattcccaggagttctttgaccatgt gaagaagctgtgggacgatgaaggagtgaaggcctgctttgagagatccaacgagtaccagc tgatcgactgtgcacaatagtaagttgcttcccaggccaggtctctctgaagcctgattgca ttcttggctgtgcccaccctatagctcagaggtttcctaaaagcagaaatcaagagtaactt tgacttcagaagttaactttctaaagaggaggttttggcaacattgtttccctgaaggtgta ttcagtaatgttgtttcaccaaaggagctggttttggtaacattgtttcactgaaggagatg tgttcagtaacattgtttcacgagcgaaccagagtaagtttctcttttgtcctcttaagttct tttcatttttttttcctatctgcacagagttaaagacatatctactaattataatgtacccaa atattcactattatattttaatttttaagcaagcctctccaaattaagttttttaatatcaa aatgtgcaggactggggagatggcttgctatctataaaagcactggatgcttttccagagga cctgagttcagttcccagcactcagctcacaaccacctataactccagctccaggggatctg gtgccctctgctgacctccttggcacacacaactaattaaaattttaaaaagtctgtttcct attgtttccctgctgcattttggagttcagcagaatgaagcttacattggtctcgggcaggt tcttactccttcgctgctgttttggagtggttccagatttgaaccagcagtgacttcagtgt ccacatactcctggtcatacaactgccctgtaaactagggtacaaagcttcaacagcaactt ttgttatcttgctgtttcctgtgggatgaatttgtgttccaggagaatgatgctcttatatt gtctgagatcaagtacaggccagtgttggatcccacccatagaatccaggtaatttgctttg tgaagaggtttaacctgtcttctctggagctttttgggataaatgagtgtggtgtgagttcc ttccgtgatggttctcaagtaataggacaacattggttgattcccttgcaaagtaaaaactg caaattaacaatccctgtgttaagacccccttcatcccttgagatgcagaaacaagcgaaac ttgctagcctggggcccagtttcaactatactgctattcataccacgaccaatgataacatc agttacctgtttaaatgccttctggggtttggtagaacataactctatagtgtcatcattta atgagttaattctaagtgcactggaactttctctgtgaaggtgaaactcacaataaagctgt tgttgtataagaagaataaataattatttcagg

Figure 19

Mouse XLGolf Protein

MGLCYSLRPLLFGSPEDTPCAASEPCAEDAQPSAAPAPASIPAPAPVGTLLRRGGGRIVANARPPGELQS' RRRQEQLRAEEREAAKEARKVSRGIDRMLREQKRDLQQTHRLLLLGAGESGKSTIVKQMRILHVNGFN PEEKKQKILDIRKNVKDAIVTIVSAMSTIIPPVPLANPENQFRSDYIKSIAPITDFEYSQEFFDHVKKLWDD EGVKACFERSNEYQLIDCAQ